

Appl. No. 10/026,140
Amdt. dated March 5, 2004
Reply to Office action of October 7, 2003

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IN THE SPECIFICATION:

Please amend the paragraph at page 18, lines 26-31 as follows:

A Basic BLASTN search (<http://www.ncbi.nlm.nih.gov/BLAST>) of the non-redundant nucleic acid sequence database was conducted on October 30, 2001, with the *bgl5* gene sequence presented in Figure 1 (SEQ ID NO:1), indicated that the only sequence producing significant alignments (i.e. with an E value of 10^{-5} or less) was GenBank Accession Number AB003109 (*bgl4* gene for beta-glucosidase of *Humicola grisea* var. *thermoidea*; E value 3×10^{-5}).

Please amend the paragraph at page 28, lines 28-32 as follows:

Preferred culture conditions for a given filamentous fungus may be found in the scientific literature and/or from the source of the fungi such as the American Type Culture Collection (ATCC; <http://www.atcc.org/>). After fungal growth has been established, the cells are exposed to conditions effective to cause or permit the over expression of BGL5.

Please amend the paragraph at page 37, lines 20-24 as follows:

Exemplary computer programs which can be used to determine identity between two sequences include, but are not limited to, the suite of BLAST programs, e.g., BLASTN, BLASTX, and TBLASTX, BLASTP and TBLASTN, publicly available on the Internet at <http://www.ncbi.nlm.nih.gov/BLAST/>. See also, Altschul, *et al.*, 1990 and Altschul, *et al.*, 1997.